



RB-125-SEQ SEQUENCE LISTING

Sequence File Name: RB125seq.txt

<110> Horwath, K. L. and Easton, C. M.

<120> Nucleic Acid Sequences Encoding Type III Tenebrio Antifreeze Proteins and Method for Assaying Activity.

<130> RB-125-SEQ

<140> 09/876,796

<141> 2001-06-07

<150> 60/210,446

<151> 2000-06-08

<160> 48

<170> Microsoft Word

<210> 1

<211> 19

<212> PRT

<213> Tenebrio molitor

<223> N-terminal sequence of protein Tm 12.86

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1 5 10 15

Gln Gln Val

19

<210> 2

<211> 576

<212> DNA

<213> Tenebrio molitor

<223> Non-his-tagged, signal plus, Tm 13.17

<400> 2

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Met Lys Leu Leu

-15

tgt tgt cta atc tcc ctc att ctg ttg gtc,aca gtt cag gcc ctg 91

Cys Cys Leu Ile Ser Leu Ile Leu Leu Val Thr Val Gln Ala Leu

-10 -5 1

acc gag gca caa att gag aaa ctg aac aag atc agc aaa aaa tgt 136

Thr Glu Ala Gln Ile Glu Lys Leu Asn Lys Ile Ser Lys Lys Cys

5 10 15

caa aat gaa agt gga gtg tcg caa gag atc ata acc aaa gct cgc 181

Gln Asn Glu Ser Gly Val Ser Gln Glu Ile Ile Thr Lys Ala Arg

20 25 30

aac ggt gac tgg gag gac gat cct aaa ctg aaa cgc caa gtt ttt 226
 Asn Gly Asp Trp Glu Asp Asp Pro Lys Leu Lys Arg Gln Val Phe
 35 40 45
 tgc gtg gcc agg aac gcc ggt ctg gcc acg gaa tcg gga gag gtg 271
 Cys Val Ala Arg Asn Ala Gly Leu Ala Thr Glu Ser Gly Glu Val
 50 55 60
 gtg gtc gac gtg ttg agg gag aag gtg agg aag gtc act gac aac 316
 Val Val Asp Val Leu Arg Glu Lys Val Arg Lys Val Thr Asp Asn
 65 70 75
 gac gaa gaa act gag aaa atc atc aat aag tgc gcc gtc aag aga 361
 Asp Glu Glu Thr Glu Lys Ile Ile Asn Lys Cys Ala Val Lys Arg
 80 85 90
 gat act gtt gaa gag acg gtg ttc aat act ttc aaa tgt gtc atg 406
 Asp Thr Val Glu Glu Thr Val Phe Asn Thr Phe Lys Cys Val Met
 95 100 105
 aaa aac aag cca aag ttc tca cca gtt gat tga accaccacga 449
 Lys Asn Lys Pro Lys Phe Ser Pro Val Asp
 110 115
 ctagtagatg gttcaaattgg tgtgctttac atataaāaat aaagtgtttc 499
 tgatgtaaaa aaaaaaaaaa aaaaaaaaaa aactcgagag tattctagag 549
 cggccgcggg cccatcgttt tccaccc 576

<210> 3

<211> 134

<212> PRT

<213> Tenebrio molitor

<223> Precursor Protein for Tm 13.17

<400> 3

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 1 5 10
 Lys Cys Gln Asn Glu Ser Gly Val Ser Gln Glu Ile Ile Thr Lys Ala
 15 20 25 30
 Arg Asn Gly Asp Trp Glu Asp Asp Pro Lys Leu Lys Arg Gln Val Phe
 35 40 45
 Cys Val Ala Arg Asn Ala Gly Leu Ala Thr Glu Ser Gly Glu Val Val
 50 55 60
 Val Asp Val Leu Arg Glu Lys Val Arg Lys Val Thr Asp Asn Asp Glu
 65 70 75

Glu Thr Glu Lys Ile Ile Asn Lys Cys Ala Val Lys Arg Asp Thr Val
 80 85 90

Glu Glu Thr Val Phe Asn Thr Phe Lys Cys Val Met Lys Asn Lys Pro
 95 100 105 110

Lys Phe Ser Pro Val Asp
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<210> 4
 <211> 116
 <212> PRT
 <213> Tenebrio molitor

<223> Mature Protein for Tm 13.17

<400> 4
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 1 5 10 15

Gln Asn Glu Ser Gly Val Ser Gln Glu Ile Ile Thr Lys Ala Arg Asn
 20 25 30

Gly Asp Trp Glu Asp Asp Pro Lys Leu Lys Arg Gln Val Phe Cys Val
 35 40 45

Ala Arg Asn Ala Gly Leu Ala Thr Glu Ser Gly Glu Val Val Val Asp
 50 55 60

Val Leu Arg Glu Lys Val Arg Lys Val Thr Asp Asn Asp Glu Glu Thr
 65 70 75 80

Glu Lys Ile Ile Asn Lys Cys Ala Val Lys Arg Asp Thr Val Glu Glu
 85 90 95

Thr Val Phe Asn Thr Phe Lys Cys Val Met Lys Asn Lys Pro Lys Phe
 100 105 110

Ser Pro Val Asp
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<210> 5
 <211> 481
 <212> DNA
 <213> Tenebrio molitor

<223> Non-His-tagged, Signal plus, Clone 2.2

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 Met Lys Leu Leu Leu Cys Phe Ala Phe Ala Ala
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 Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys

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Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val Ser			
10	15	20	
caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gat gat			181
Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp			
25	30	35	
ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act gga			226
Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly			
40	45	50	
gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa gcc			271
Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala			
55	60	65	
aag ctg aag cat gtg gcc agc gac gaa gag gtg gac aag atc gtg			316
Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val			
70	75	80	
cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct tat			361
Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr			
85	90	95	
gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct cct			406
Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro			
100	105	110	
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Ile Asp			
115			
tcgttatgta aaaaaaaaaa aaaaaa			481
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<213> Tenebrio molitor			
<223> Non-His-tagged, Signal plus, Clone 2.3			
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atc gtc atc gga gct cag gct ctc acc gac gaa cag ata cag aaa			91
Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys			
-5	1	5	
agg aac aag atc agc aaa gaa tgc cag cag gtg tcc gga gtg tcc			136
Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val Ser			
10	15	20	

caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gac gat 181
 Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp
 25 30 35

 ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act gga 226
 Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly
 40 45 50

 gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa gcc 271
 Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala
 55 60 65

 aag ctg aag cat gtg gcc agc gac gaa gaa gtg gac aag atc gtg 316
 Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val
 70 75 80

 cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct tat 361
 Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr
 85 90 95

 gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct cct 406
 Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro
 100 105 110
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 Ile Asp
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 tcgttatgaa aaaaaaaaaa aaaaaaa 482

<210> 7
 <211> 133
 <212> PRT
 <213> Tenebrio molitor

<223> Precursor Protein for Tm 12.84, Clones 2.2, 2.3, and 7.5

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Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val
 15 20 25 30

Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys His Val Leu
 35 40 45

Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala Gly Asp Thr Asn
 50 55 60

Val Glu Val Leu Lys Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu
 65 70 75

Val Asp Lys Ile Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu

80 85 90

Glu Thr Ala Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp
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Phe Ser Pro Ile Asp
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<210> 8
 <211> 115
 <212> PRT
 <213> Tenebrio molitor

<223> Mature Protein for Tm 12.84, Clones 2.2, 2.3, and 7.5

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Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val Arg Thr
 20 25 30

Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys His Val Leu Cys Phe
 35 40 45

Ser Lys Lys Thr Gly Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu
 50 55 60

Val Leu Lys Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp
 65 70 75 80

Lys Ile Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr
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Ala Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser
 100 105 110

Pro Ile Asp
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<210> 9
 <211> 481
 <212> DNA
 <213> Tenebrio molitor

<223> Non-His-tagged, Signal plus, Clone 3.4

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 -15 -10

atc gtc atc gga gct cag gct ctc acc gac gaa cag ata cag aaa 91
 Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys
 -5 1 5


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agg aac aag atc agc aaa gaa tgc cag cag gtg tcc gga gtg tcc 136
Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val Ser
10 15 20

caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gac gat 181
Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp
25 30 35

ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act gga 226
Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly
40 45 50

gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa gcc 271
Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala
55 60 65

aag ctg aag cat gtg gcc agc gac gaa gag gtg gac aag atc gtg 316
Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val
70 75 80

cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct tat 361
Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr
85 90 95

gac acc ttc aag gtt att tac gac agt aaa cct gat ttc tct cct 406
Asp Thr Phe Lys Val Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro
100 105 110

att gat taa ttgttttgta tttgactgaa ttttgacaat aaaggtacta 455
Ile Asp
115

tcgttatgta aaaaaaaaaa aaaaaa 481

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<210> 10
<211> 133
<212> PRT
<213> Tenebrio molitor

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<223> Precursor Protein for Clone 3.4

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1 5 10

Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val
15 20 25 30

Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys His Val Leu
35 40 45

Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala Gly Asp Thr Asn

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50 55 60
 Val Glu Val Leu Lys Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu
 65 70 75
 Val Asp Lys Ile Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu
 80 85 90
 Glu Thr Ala Tyr Asp Thr Phe Lys Val Ile Tyr Asp Ser Lys Pro Asp
 95 100 105 110
 Phe Ser Pro Ile Asp
 115

<210> 11
 <211> 115
 <212> PRT
 <213> Tenebrio molitor

<223> Mature Protein for Clone 3.4

<400> 11
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 20 25 30
 Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys His Val Leu Cys Phe
 35 40 45
 Ser Lys Lys Thr Gly Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu
 50 55 60
 Val Leu Lys Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp
 65 70 75 80
 Lys Ile Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr
 85 90 95
 Ala Tyr Asp Thr Phe Lys Val Ile Tyr Asp Ser Lys Pro Asp Phe Ser
 100 105 110
 Pro Ile Asp
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<210> 12
 <211> 482
 <212> DNA
 <213> Tenebrio molitor

<223> Non-His-tagged, Signal plus, Clone 3.9

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 -15 -10

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 Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys
 -5 1 5

agg aac aag atc agc aaa gaa tgc cag cag gag tcc gga gtg tcc 136
 Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Glu Ser Gly Val Ser
 10 15 20

caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gac gat 181
 Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp
 25 30 35

ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aga act gga 226
 Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Arg Thr Gly
 40 45 50

gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa gcc 271
 Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala
 55 60 65

aag ctg aag cat gtg gcc agc gac gaa gaa gtg gac aag atc gtg 316
 Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val
 70 75 80

cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct tat 361
 Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr
 85 90 95

gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct cct 406
 Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro
 100 105 110

att gat taa ttgttttgta tttgactgaa ttttgacaat aaaggacta 455
 Ile Asp
 115

tcgttatgaa aaaaaaaaaa aaaaaaa 482

<210> 13
 <211> 133
 <212> PRT
 <213> Tenebrio molitor

<223> Precursor Protein for Clone 3.9

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 1 5 10

Glu Cys Gln Gln Glu Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val

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Arg	Thr	Gly	Val	Leu	Val	Asp	Asp	Pro	Lys	Met	Lys	Lys	His	Val	Leu
			35						40					45	
Cys	Phe	Ser	Lys	Arg	Thr	Gly	Val	Ala	Thr	Glu	Ala	Gly	Asp	Thr	Asn
			50					55					60		
Val	Glu	Val	Leu	Lys	Ala	Lys	Leu	Lys	His	Val	Ala	Ser	Asp	Glu	Glu
		65					70					75			
Val	Asp	Lys	Ile	Val	Gln	Lys	Cys	Val	Val	Lys	Lys	Ala	Thr	Pro	Glu
	80					85					90				
Glu	Thr	Ala	Tyr	Asp	Thr	Phe	Lys	Cys	Ile	Tyr	Asp	Ser	Lys	Pro	Asp
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Phe	Ser	Pro	Ile	Asp											
				115											

<210> 14
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 <212> PRT
 <213> Tenebrio molitor

<223> Mature protein for Clone 3.9

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Gln	Gln	Glu	Ser	Gly	Val	Ser	Gln	Glu	Thr	Ile	Asp	Lys	Val	Arg	Thr
			20					25					30		
Gly	Val	Leu	Val	Asp	Asp	Pro	Lys	Met	Lys	Lys	His	Val	Leu	Cys	Phe
		35					40					45			
Ser	Lys	Arg	Thr	Gly	Val	Ala	Thr	Glu	Ala	Gly	Asp	Thr	Asn	Val	Glu
	50					55					60				
Val	Leu	Lys	Ala	Lys	Leu	Lys	His	Val	Ala	Ser	Asp	Glu	Glu	Val	Asp
65					70				75					80	
Lys	Ile	Val	Gln	Lys	Cys	Val	Val	Lys	Lys	Ala	Thr	Pro	Glu	Glu	Thr
			85						90					95	
Ala	Tyr	Asp	Thr	Phe	Lys	Cys	Ile	Tyr	Asp	Ser	Lys	Pro	Asp	Phe	Ser
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Pro	Ile	Asp													
		115													

<210> 15
 <211> 481
 <212> DNA

<213> Tenebrio molitor

<223> Non-his-tagged, Signal plus, Clone 7.5

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-10

atc gtc atc gga gct cag gct ctc acc gac gaa cag ata cag aaa 91

Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys

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agg aac aag atc agc aaa gag tgc cag cag gtg tcc gga gtg tcc 136

Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val Ser

10

15

20

caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gac gat 181

Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp

25

30

35

ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act gga 226

Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly

40

45

50

gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa gcc 271

Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala

55

60

65

aag ctg aag cat gtg gcc agc gac gaa gag gtg gac aag atc gtg 316

Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val

70

75

80

cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct tat 361

Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr

85

90

95

gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct cct 406

Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro

100

105

110

att gat taa ttgttttgta ttggttgaa ttttgacaat aaaggtacta 455

Ile Asp

115

tcgttatgta aaaaaaaaaa aaaaaa 481

<210> 16

<211> 681

<212> DNA

<213> Tenebrio molitor

<223> His-tagged, Signal plus, Clone 2.2

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agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly -45 -40 -35	141
gga cag caa atg ggt cgc gga tcc gaa ttc gca cga gca aaa atg Gly Gln Gln Met Gly Arg Gly Ser Glu Phe Ala Arg Ala Lys Met -30 -25 -20	186
aaa ctc ctc ttg tgc ttt gcg ttc gcc gcc atc gtc atc gga gct Lys Leu Leu Leu Cys Phe Ala Phe Ala Ala Ile Val Ile Gly Ala -15 -10 -5	231
cag gct ctc acc gac gaa cag ata cag aaa agg aac aag atc agc Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser 1 5 10	276
aaa gaa tgc cag cag gtg tcc gga gtg tcc caa gag acg atc gac Lys Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp 15 20 25	321
aaa gtc cgc aca ggt gtc ttg gtc gat gat ccc aaa atg aag aag Lys Val Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys 30 35 40	366
cac gtc ctc tgc ttc tcg aag aaa act gga gtg gca acc gaa gcc His Val Leu Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala 45 50 55	411
gga gac acc aat gtg gag gta ctc aaa gcc aag ctg aag cat gtg Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu Lys His Val 60 65 70	456
gcc agc gac gaa gag gtg gac aag atc gtg cag aag tgc gtg gtc Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys Val Val 75 80 85	501
aag aag gcc aca cca gag gaa acg gct tat gac acc ttc aag tgt Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys Cys 90 95 100	546
att tac gac agt aaa cct gat ttc tct cct att gat taa ttgttttgta Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp 105 110 115	595
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aaaaaactcg agcaccacca ccaccaccac tga gat	681

<210> 17
 <211> 173
 <212> PRT

<213> Tenebrio molitor

<223> Precursor Protein with His-tag, Clone 2.2

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-40 -35 -30

Gly Ser Glu Phe Ala Arg Ala Lys Met Lys Leu Leu Leu Cys Phe Ala
-25 -20 -15

Phe Ala Ala Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile
-10 -5 1 5

Gln Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val
10 15 20

Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp
25 30 35

Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly Val
40 45 50

Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu
55 60 65 70

Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys
75 80 85

Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys
90 95 100

Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp
105 110 115

<210> 18

<211> 543

<212> DNA

<213> Tenebrio molitor

<223> His-tagged, signal minus, Clone 2.2

<400> 18

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-30 -25

agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt 141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly
-20 -15 -10

gga cag caa atg ggt cgc gga tcc ctc acc gac gaa cag ata cag 186
 Gly Gln Gln Met Gly Arg Gly Ser Leu Thr Asp Glu Gln Ile Gln
 -5 1 5
 aaa agg aac aag atc agc aaa gaa tgc cag-cag gtg tcc gga gtg 231
 Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val
 10 15 20
 tcc caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gat 276
 Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp
 25 30 35
 gat ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act 321
 Asp Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr
 40 45 50
 gga gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa 366
 Gly Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys
 55 60 65
 gcc aag ctg aag cat gtg gcc agc gac gaa gag gtg gac aag atc 411
 Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile
 70 75 80
 gtg cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct 456
 Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala
 85 90 95
 tat gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct 501
 Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser
 100 105 110
 cct att gat taa ctgcagcacc accaccacca ccaactgagat 543
 Pro Ile Asp
 115

<210> 19
 <211> 149
 <212> PRT
 <213> Tenebrio molitor

<223> Mature Protein with His-tag, Clone 2.2

<400> 19
 Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
 -30 -25 -20
 Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
 -15 -10 -5
 Gly Ser Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser Lys
 1 5 10
 Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val
 15 20 25 30

cac gtc ctc tgc ttc tcg aag aaa act gga gtg gca acc gaa gcc	411
His Val Leu Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala	
45 50 55	
gga gac acc aat gtg gag gta ctc aaa gcc-aag ctg aag cat gtg	456
Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu Lys His Val	
60 65 70	
gcc agc gac gaa gaa gtg gac aag atc gtg cag aag tgc gtg gtc	501
Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys Val Val	
75 80 85	
aag aag gcc aca cca gag gaa acg gct tat gac acc ttc aag tgt	546
Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys Cys	
90 95 100	
att tac gac agt aaa cct gat ttc tct cct att gat taa ttgttttgta	595
Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp	
105 110 115	
tttgactgaa ttttgacaat aaaggtacta tcgttatgaa aaaaaaaaaa	645
aaaaaaaaactc gagcaccacc accaccacca ctgagat	682

<210> 21
 <211> 173
 <212> PRT
 <213> Tenebrio molitor

<223> Precursor Protein with His-tag, Clone 2.3

<400> 21
 Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
 -55 -50 -45
 Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
 -40 -35 -30
 Gly Ser Glu Phe Ala Arg Ala Lys Met Lys Leu Leu Leu Cys Phe Ala
 -25 -20 -15
 Phe Ala Ala Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile
 -10 -5 1 5
 Gln Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val
 10 15 20
 Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp
 25 30 35
 Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly Val
 40 45 50
 Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu
 55 60 65 70

Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys
75 80 85

Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys
90 95 100

Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp
105 110 115

<210> 22
<211> 543
<212> DNA
<213> *Tenebrio molitor*

<223> His-tagged, Signal minus, Clone 2.3

<400> 22
ttgttagcgg atggaattcc ctcgtagggg ataattttgt ttactttaag 50
aaggagatat acc atg ggc agc agc cat cat cat cat cat cac agc 96
Met Gly Ser Ser His His His His His His Ser
-30 -25
agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt 141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly
-20 -15 -10
gga cag caa atg ggt cgc gga tcc ctc acc gac gaa cag ata cag 186
Gly Gln Gln Met Gly Arg Gly Ser Leu Thr Asp Glu Gln Ile Gln
-5 1 5
aaa agg aac aag atc agc aaa gaa tgc cag cag gtg tcc gga gtg 231
Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val
10 15 20
tcc caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gat 276
Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp
25 30 35
gat ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act 321
Asp Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr
40 45 50
gga gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa 366
Gly Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys
55 60 65
gcc aag ctg aag cat gtg gcc agc gac gaa gaa gtg gac aag atc 411
Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile
70 75 80
gtg cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct 456
Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala
85 90 95
tat gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct 501

Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser
 100 105 110

cct att gat taa ctcgagcacc accaccacca ccactgagat 543
 Pro Ile Asp
 115

<210> 23
 <211> 149
 <212> PRT
 <213> Tenebrio molitor

<223> Mature Protein with His-tag, Clone 2.3

<400> 23
 Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
 -30 -25 -20

Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
 -15 -10 -5

Gly Ser Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser Lys
 1 5 10

Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val
 15 20 25 30

Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys His Val Leu
 35 40 45

Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala Gly Asp Thr Asn
 50 55 60

Val Glu Val Leu Lys Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu
 65 70 75

Val Asp Lys Ile Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu
 80 85 90

Glu Thr Ala Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp
 95 100 105 110

Phe Ser Pro Ile Asp
 115

<210> 24
 <211> 776
 <212> DNA
 <213> Tenebrio molitor

<223> His-tagged, Signal plus, Tm 13.17

<400> 24
 ttgttagcgg atggaattcc ctcgtagggg ataattttgt ttactttaag 50

aaggagatat acc atg ggc agc agc cat cat cat cat cat cac agc	96
Met Gly Ser Ser His His His His His His Ser	
-65 -60 -55	
agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt	141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly	
-50 -45 -40	
gga cag caa atg ggt cgc gga tcc gaa ttc tgg atc caa aga att	186
Gly Gln Gln Met Gly Arg Gly Ser Glu Phe Trp Ile Gln Arg Ile	
-35 -30 -25	
cgg cac gag act act aag atg aag ttg ctc tgt tgt cta atc tcc	231
Arg His Glu Thr Thr Lys Met Lys Leu Leu Cys Cys Leu Ile Ser	
-20 -15 -10	
ctc att ctg ttg gtc aca gtt cag gcc ctg acc gag gca caa att	276
Leu Ile Leu Leu Val Thr Val Gln Ala Leu Thr Glu Ala Gln Ile	
-5 1 5	
gag aaa ctg aac aag atc agc aaa aaa tgt caa aat gaa agt gga	321
Glu Lys Leu Asn Lys Ile Ser Lys Lys Cys Gln Asn Glu Ser Gly	
10 15 20	
gtg tcg caa gag atc ata acc aaa gct cgc aac ggt gac tgg gag	366
Val Ser Gln Glu Ile Ile Thr Lys Ala Arg Asn Gly Asp Trp Glu	
25 30 35	
gac gat cct aaa ctg aaa cgc caa gtt ttt tgc gtg gcc agg aac	411
Asp Asp Pro Lys Leu Lys Arg Gln Val Phe Cys Val Ala Arg Asn	
40 45 50	
gcc ggt ctg gcc acg gaa tcg gga gag gtg gtg gtc gac gtg ttg	456
Ala Gly Leu Ala Thr Glu Ser Gly Glu Val Val Val Asp Val Leu	
55 60 65	
agg gag aag gtg agg aag gtc act gac aac gac gaa gaa act gag	501
Arg Glu Lys Val Arg Lys Val Thr Asp Asn Asp Glu Glu Thr Glu	
70 75 80	
aaa atc atc aat aag tgc gcc gtc aag aga gat act gtt gaa gag	546
Lys Ile Ile Asn Lys Cys Ala Val Lys Arg Asp Thr Val Glu Glu	
85 90 95	
acg gtg ttc aat act ttc aaa tgt gtc atg aaa aac aag cca aag	591
Thr Val Phe Asn Thr Phe Lys Cys Val Met Lys Asn Lys Pro Lys	
100 105 110	
ttc tca cca gtt gat tga accaccacga ctagtagatg gttcaaatgg	639
Phe Ser Pro Val Asp	
115	
tgtgctttac atataaaaat aaagtgtttc tgatgtaaaa aaaaaaaaaa	689
aaaaaaaaaa aactcgagag tattctagag cggccgcggg cccatcgttt	739

tccacccctc gagcaccacc accaccacca ctgagat

776

<210> 25
<211> 174
<212> PRT
<213> Tenebrio molitor

<223> Precursor Protein with His-tag, Tm 13.17

<400> 25
Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
 -55 -50 -45

Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
 -40 -35 -30

Gly Ser Glu Phe Ala Arg Ala Lys Met Lys Leu Leu Cys Cys Leu Ile
 -25 -20 -15

Ser Leu Ile Leu Leu Val Thr Val Gln Ala Leu Thr Glu Ala Gln Ile
 -10 -5 1 5

Glu Lys Leu Asn Lys Ile Ser Lys Lys Cys Gln Asn Glu Ser Gly Val
 10 15 20

Ser Gln Glu Ile Ile Thr Lys Ala Arg Asn Gly Asp Trp Glu Asp Asp
 25 30 35

Pro Lys Leu Lys Arg Gln Val Phe Cys Val Ala Arg Asn Ala Gly Leu
 40 45 50

Ala Thr Glu Ser Gly Glu Val Val Val Asp Val Leu Arg Glu Lys Val
55 60 65 70

Arg Lys Val Thr Asp Asn Asp Glu Glu Thr Glu Lys Ile Ile Asn Lys
 75 80 85

Cys Ala Val Lys Arg Asp Thr Val Glu Glu Thr Val Phe Asn Thr Phe
 90 95 100

Lys Cys Val Met Lys Asn Lys Pro Lys Phe Ser Pro Val Asp
 105 110 115

<210> 26
<211> 543
<212> DNA
<213> Tenebrio molitor

<223> His-tagged, Signal minus, Tm 13.17

<400> 26
ttgttagcgg atggaattcc ctcgtagggg ataattttgt ttactttaag 50

aaggagatat acc atg ggc agc agc cat cat cat cat cat cac agc 96
 Met Gly Ser Ser His His His His His His Ser

-30

-25

agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt 141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly
-20 -15 -10

gga cag caa atg ggt cgc ggc ctg acc gag gca caa att gag aaa 186
Gly Gln Gln Met Gly Arg Gly Leu Thr Glu Ala Gln Ile Glu Lys
-5 1 5

ctg aac aag atc agc aaa aaa tgt caa aat gaa agt gga gtg tcg 231
Leu Asn Lys Ile Ser Lys Lys Cys Gln Asn Glu Ser Gly Val Ser
10 15 20

caa gag atc ata acc aaa gct cgc aac ggt gac tgg gag gac gat 276
Gln Glu Ile Ile Thr Lys Ala Arg Asn Gly Asp Trp Glu Asp Asp
25 30 35

cct aaa ctg aaa cgc caa gtt ttt tgc gtg gcc agg aac gcc ggt 321
Pro Lys Leu Lys Arg Gln Val Phe Cys Val Ala Arg Asn Ala Gly
40 45 50

ctg gcc acg gaa tcg gga gag gtg gtg gtc gac gtg ttg agg gag 366
Leu Ala Thr Glu Ser Gly Glu Val Val Val Asp Val Leu Arg Glu
55 60 65

aag gtg agg aag gtc act gac aac gac gaa gaa act gag aaa atc 411
Lys Val Arg Lys Val Thr Asp Asn Asp Glu Glu Thr Glu Lys Ile
70 75 80

atc aat aag tgc gcc gtc aag aga gat act gtt gaa gag acg gtg 456
Ile Asn Lys Cys Ala Val Lys Arg Asp Thr Val Glu Glu Thr Val
85 90 95

ttc aat act ttc aaa tgt gtc atg aaa aac aag cca aag ttc tca 501
Phe Asn Thr Phe Lys Cys Val Met Lys Asn Lys Pro Lys Phe Ser
100 105 110

cca gtt gat tga ctgcagcacc accaccacca ccactgagat 543
Pro Val Asp
115

<210> 27
<211> 149
<212> PRT
<213> Tenebrio molitor

<223> Mature Protein with His-tag, Tm 13.17

<400> 27
Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
-30 -25 -20

Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
-15 -10 -5

Gly Leu Thr Glu Ala Gln Ile Glu Lys Leu Asn Lys Ile Ser Lys Lys
 1 5 10 15
 Cys Gln Asn Glu Ser Gly Val Ser Gln Glu Ile Ile Thr Lys Ala Arg
 20 25 30
 Asn Gly Asp Trp Glu Asp Asp Pro Lys Leu Lys Arg Gln Val Phe Cys
 35 40 45
 Val Ala Arg Asn Ala Gly Leu Ala Thr Glu Ser Gly Glu Val Val Val
 50 55 60
 Asp Val Leu Arg Glu Lys Val Arg Lys Val Thr Asp Asn Asp Glu Glu
 65 70 75
 Thr Glu Lys Ile Ile Asn Lys Cys Ala Val Lys Arg Asp Thr Val Glu
 80 85 90 95
 Glu Thr Val Phe Asn Thr Phe Lys Cys Val Met Lys Asn Lys Pro Lys
 100 105 110
 Phe Ser Pro Val Asp
 115

<210> 28
 <211> 681
 <212> DNA
 <213> *Tenebrio molitor*

<223> His-tagged, Signal plus, Clone 3.4

<400> 28
 ttgttagcgg atggaattcc ctcgtagggg ataattttgt ttactttaag 50
 aaggagatat acc atg ggc agc agc cat cat cat cat cat cac agc 96
 Met Gly Ser Ser His His His His His His Ser
 -55 -50
 agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt 141
 Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly
 -45 -40 -35
 gga cag caa atg ggt cgc gga tcc gaa ttc gca cga gca aaa atg 186
 Gly Gln Gln Met Gly Arg Gly Ser Glu Phe Ala Arg Ala Lys Met
 -30 -25 -20
 aaa ctc ctc ttg tgc ttt gct ttc gcc gcc atc gtc atc gga gct 231
 Lys Leu Leu Leu Cys Phe Ala Phe Ala Ala Ile Val Ile Gly Ala
 -15 -10 -5
 cag gct ctc acc gac gaa cag ata cag aaa agg aac aag atc agc 276
 Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser
 1 5 10
 aaa gaa tgc cag cag gtg tcc gga gtg tcc caa gag acg atc gac 321
 Lys Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp

15	20	25	
aaa gtc cgc aca ggt gtc ttg gtc gac gat ccc	aaa atg aag aag	366	
Lys Val Arg Thr Gly Val Leu Val Asp Asp Pro	Lys Met Lys Lys		
30	40		
cac gtc ctc tgc ttc tcg aag aaa act gga gtg gca acc gaa gcc	411		
His Val Leu Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala			
45	55		
gga gac acc aat gtg gag gta ctc aaa gcc aag ctg aag cat gtg	456		
Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu Lys His Val			
60	70		
gcc agc gac gaa gag gtg gac aag atc gtg cag aag tgc gtg gtc	501		
Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys Val Val			
75	85		
aag aag gcc aca cca gag gaa acg gct tat gac acc ttc aag gtt	546		
Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys Val			
90	100		
att tac gac agt aaa cct gat ttc tct cct att gat taa ttgttttgta	595		
Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp			
105	115		
tttgactgaa ttttgacaat aaagggtacta tcgttatgta aaaaaaaaaa	645		
aaaaaactcg agcaccacca ccaccaccac tgagat	681		
<210> 29			
<211> 173			
<212> PRT			
<213> Tenebrio molitor			
<223> Precursor protein with His-tag, Clone 3.4			
<400> 29			
Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro			
-55	-50	-45	
Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg			
-40	-35	-30	
Gly Ser Glu Phe Ala Arg Ala Lys Met Lys Leu Leu Leu Cys Phe Ala			
-25	-20	-15	
Phe Ala Ala Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile			
-10	-5	1	
		5	
Gln Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val			
10	15	20	
Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp			
25	30	35	

Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly Val
40 45 50

Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu
55 60 65 70

Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys
75 80 85

Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys
90 95 100

Val Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp
105 110 115

<210> 30
<211> 543
<212> DNA
<213> Tenebrio molitor

<223> His-tagged, Signal minus, Clone 3.4

<400> 30
ttgtagcgg atggaattcc ctgtagggg ataattttgt ttactttaag 50

aaggagatat acc atg ggc agc agc cat cat cat cat cac agc 96
Met Gly Ser Ser His His His His His Ser
-30 -25

agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt 141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly
-20 -15 -10

gga cag caa atg ggt cgc gga tcc ctc acc gac gaa cag ata cag 186
Gly Gln Gln Met Gly Arg Gly Ser Leu Thr Asp Glu Gln Ile Gln
-5 1 5

aaa agg aac aag atc agc aaa gaa tgc cag cag gtg tcc gga gtg 231
Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val
10 15 20

tcc caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gac 276
Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp
25 30 35

gat ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act 321
Asp Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr
40 45 50

gga gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa 366
Gly Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys
55 60 65

gcc aag ctg aag cat gtg gcc agc gac gaa gag gtg gac aag atc 411
Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile
70 75 80

gtg cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct 456
 Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala
 85 90 95

tat gac acc ttc aag gtt att tac gac agt aaa cct gat ttc tct 501
 Tyr Asp Thr Phe Lys Val Ile Tyr Asp Ser Lys Pro Asp Phe Ser
 100 105 110

cct att gat taa ctcgagcacc accaccacca ccactgagat 543
 Pro Ile Asp
 115

<210> 31
 <211> 149
 <212> PRT
 <213> Tenebrio molitor

<223> Mature Protein with His-tag, Clone 3.4

<400> 31
 Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
 -30 -25 -20

Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
 -15 -10 -5

Gly Ser Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser Lys
 1 5 10

Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val
 15 20 25 30

Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys His Val Leu
 35 40 45

Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala Gly Asp Thr Asn
 50 55 60

Val Glu Val Leu Lys Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu
 65 70 75

Val Asp Lys Ile Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu
 80 85 90

Glu Thr Ala Tyr Asp Thr Phe Lys Val Ile Tyr Asp Ser Lys Pro Asp
 95 100 105 110

Phe Ser Pro Ile Asp
 115

<210> 32
 <211> 682
 <212> DNA
 <213> Tenebrio molitor

<223> His-tagged, Signal plus, Clone 3.9

<400> 32

ttgttagcgg atggaattcc ctcgtagggg ataattttgt ttactttaag	50
aaggagatat acc atg ggc agc agc cat cat cat cat cat cac agc	96
Met Gly Ser Ser His His His His His His Ser	
-55 -50	
agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt	141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly	
-45 -40 -35	
gga cag caa atg ggt cgc gga tcc gaa ttc gca cga gca aaa atg	186
Gly Gln Gln Met Gly Arg Gly Ser Glu Phe Ala Arg Ala Lys Met	
-30 -25 -20	
aaa ctc ctc ttg tgc ttt gct ttc gcc gcc atc gtc atc gga gct	231
Lys Leu Leu Leu Cys Phe Ala Phe Ala Ala Ile Val Ile Gly Ala	
-15 -10 -5	
cag gct ctc acc gat gaa cag ata cag aaa agg aac aag atc agc	276
Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser	
1 5 10	
aaa gaa tgc cag cag gag tcc gga gtg tcc caa gag acg atc gac	321
Lys Glu Cys Gln Gln Glu Ser Gly Val Ser Gln Glu Thr Ile Asp	
15 20 25	
aaa gtc cgc aca ggt gtc ttg gtc gac gat ccc aaa atg aag aag	366
Lys Val Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys	
30 35 40	
cac gtc ctc tgc ttc tcg aag aga act gga gtg gca acc gaa gcc	411
His Val Leu Cys Phe Ser Lys Arg Thr Gly Val Ala Thr Glu Ala	
45 50 55	
gga gac acc aat gtg gag gta ctc aaa gcc aag ctg aag cat gtg	456
Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu Lys His Val	
60 65 70	
gcc agc gac gaa gaa gtg gac aag atc gtg cag aag tgc gtg gtc	501
Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys Val Val	
75 80 85	
aag aag gcc aca cca gag gaa acg gct tat gac acc ttc aag tgt	546
Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys Cys	
90 95 100	
att tac gac agt aaa cct gat ttc tct cct att gat taa ttgttttgta	595
Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp	
105 110 115	
tttgactgaa ttttgacaat aaaggtagta tcgttatgaa aaaaaaaaaa	645
aaaaaaaaactc gagcaccacc accaccacca ctgagat	682

<210> 33
 <211> 173
 <212> PRT
 <213> Tenebrio molitor

<223> Precursor Protein with His-tag, Clone 3.9

<400> 33
 Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
 -55 -50 -45
 Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
 -40 -35 -30
 Gly Ser Glu Phe Ala Arg Ala Lys Met Lys Leu Leu Leu Cys Phe Ala
 -25 -20 -15
 Phe Ala Ala Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile
 -10 -5 1 5
 Gln Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Glu Ser Gly Val
 10 15 20
 Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp
 25 30 35
 Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Arg Thr Gly Val
 40 45 50
 Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu
 55 60 65 70
 Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys
 75 80 85
 Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys
 90 95 100
 Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp
 105 110 115

<210> 34
 <211> 543
 <212> DNA
 <213> Tenebrio molitor

<223> His-tagged, Signal minus, Clone 3.9

<400> 34
 ttgttagcgg atggaattcc ctgtagggg ataattttgt ttactttaag 50
 aaggagatat acc atg ggc agc agc cat cat cat cat cac agc 96
 Met Gly Ser Ser His His His His His His Ser
 -30 -25

agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt 141
 Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly
 -20 -15 -10

gga cag caa atg ggt cgc gga tcc ctc acc gat gaa cag ata cag 186
 Gly Gln Gln Met Gly Arg Gly Ser Leu Thr Asp Glu Gln Ile Gln
 -5 1 5

aaa agg aac aag atc agc aaa gaa tgc cag cag gag tcc gga gtg 231
 Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Glu Ser Gly Val
 10 15 20

tcc caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gac 276
 Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp
 25 30 35

gat ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aga act 321
 Asp Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Arg Thr
 40 45 50

gga gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa 366
 Gly Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys
 55 60 65

gcc aag ctg aag cat gtg gcc agc gac gaa gaa gtg gac aag atc 411
 Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile
 70 75 80

gtg cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct 456
 Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala
 85 90 95

tat gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct 501
 Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser
 100 105 110

cct att gat taa ctgcgagcacc accaccacca ccaactgagat 543
 Pro Ile Asp
 115

<210> 35
 <211> 149
 <212> PRT
 <213> Tenebrio molitor

<223> Mature Protein with His-tag, Clone 3.9

<400> 35
 Met Gly Ser Ser His His His His His Ser Ser Gly Leu Val Pro
 -30 -25 -20

Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
 -15 -10 -5

Gly Ser Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser Lys

1					5					10					
Glu	Cys	Gln	Gln	Glu	Ser	Gly	Val	Ser	Gln	Glu	Thr	Ile	Asp	Lys	Val
15					20					25					30
Arg	Thr	Gly	Val	Leu	Val	Asp	Asp	Pro	Lys	Met	Lys	Lys	His	Val	Leu
			35					40					45		
Cys	Phe	Ser	Lys	Arg	Thr	Gly	Val	Ala	Thr	Glu	Ala	Gly	Asp	Thr	Asn
			50					55					60		
Val	Glu	Val	Leu	Lys	Ala	Lys	Leu	Lys	His	Val	Ala	Ser	Asp	Glu	Glu
	65					70					75				
Val	Asp	Lys	Ile	Val	Gln	Lys	Cys	Val	Val	Lys	Lys	Ala	Thr	Pro	Glu
	80					85					90				
Glu	Thr	Ala	Tyr	Asp	Thr	Phe	Lys	Cys	Ile	Tyr	Asp	Ser	Lys	Pro	Asp
95					100					105					110
Phe	Ser	Pro	Ile	Asp											
			115												

<210> 36
 <211> 681
 <212> DNA
 <213> Tenebrio molitor

<223> His-tagged, Signal plus, Clone 7.5

<400> 36	
ttgttagcgg atggaattcc ctcgtagggg ataattttgt ttactttaag	50
aaggagatat acc atg ggc agc agc cat cat cat cat cat cac agc	96
Met Gly Ser Ser His His His His His His Ser	
-55 -50	
agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt	141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly	
-45 -40 -35	
gga cag caa atg ggt cgc gga tcc gaa ttc gca cga gca aaa atg	186
Gly Gln Gln Met Gly Arg Gly Ser Glu Phe Ala Arg Ala Lys Met	
-30 -25 -20	
aaa ctc ctc ttg tgc ttt gcg ttc gcc gcc atc gtc atc gga gct	231
Lys Leu Leu Leu Cys Phe Ala Phe Ala Ala Ile Val Ile Gly Ala	
-15 -10 -5	
cag gct ctc acc gac gaa cag ata cag aaa agg aac aag atc agc	276
Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser	
1 5 10	
aaa gag tgc cag cag gtg tcc gga gtg tcc caa gag acg atc gac	321
Lys Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp	
15 20 25	

aaa gtc cgc aca ggt gtc ttg gtc gac gat ccc aaa atg aag aag	366
Lys Val Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys	
30 35 40	

cac gtc ctc tgc ttc tcg aag aaa act gga gtg gca acc gaa gcc	411
His Val Leu Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala	
45 50 55	

gga gac acc aat gtg gag gta ctc aaa gcc aag ctg aag cat gtg	456
Gly Asp Thr Asn Val Glu Val Leu Lys Ala Lys Leu Lys His Val	
60 65 70	

gcc agc gac gaa gag gtg gac aag atc gtg cag aag tgc gtg gtc	501
Ala Ser Asp Glu Glu Val Asp Lys Ile Val Gln Lys Cys Val Val	
75 80 85	

aag aag gcc aca cca gag gaa acg gct tat gac acc ttc aag tgt	546
Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr Asp Thr Phe Lys Cys	
90 95 100	

att tac gac agt aaa cct gat ttc tct cct att gat taa ttgttttgta	595
Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro Ile Asp	
105 110 115	

tttggctgaa ttttgacaat aaaggtacta tcgttatgta aaaaaaaaaa	645
--	-----

aaaaaactcg agcaccacca ccaccaccac tgagat	681
---	-----

<210> 37
 <211> 173
 <212> PRT
 <213> Tenebrio molitor

<223> Precursor Protein with His-tag, Clone 7.5

<400> 37	
Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro	
-55 -50 -45	
Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg	
-40 -35 -30	
Gly Ser Glu Phe Ala Arg Ala Lys Met Lys Leu Leu Leu Cys Phe Ala	
-25 -20 -15	
Phe Ala Ala Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile	
-10 -5 1 5	
Gln Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val	
10 15 20	
Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp	

25					30					35					
Pro	Lys	Met	Lys	Lys	His	Val	Leu	Cys	Phe	Ser	Lys	Lys	Thr	Gly	Val
40						45					50				
Ala	Thr	Glu	Ala	Gly	Asp	Thr	Asn	Val	Glu	Val	Leu	Lys	Ala	Lys	Leu
55					60					65					70
Lys	His	Val	Ala	Ser	Asp	Glu	Glu	Val	Asp	Lys	Ile	Val	Gln	Lys	Cys
				75					80					85	
Val	Val	Lys	Lys	Ala	Thr	Pro	Glu	Glu	Thr	Ala	Tyr	Asp	Thr	Phe	Lys
			90					95					100		
Cys	Ile	Tyr	Asp	Ser	Lys	Pro	Asp	Phe	Ser	Pro	Ile	Asp			
	105						110					115			

<210> 38
 <211> 543
 <212> DNA
 <213> Tenebrio molitor

<223> His-tagged, Signal minus, Clone 7.5

<400> 38	
ttggttagcgg atggaattcc ctgtagggg ataattttgt ttactttaag	50
aaggagatat acc atg ggc agc agc cat cat cat cat cac agc	96
Met Gly Ser Ser His His His His His Ser	
-30 -25	
agc ggc ctg gtg ccg cgc ggc agc cat atg gct agc atg act ggt	141
Ser Gly Leu Val Pro Arg Gly Ser His Met Ala Ser Met Thr Gly	
-20 -15 -10	
gga cag caa atg ggt cgc gga tcc ctc acc gac gaa cag ata cag	186
Gly Gln Gln Met Gly Arg Gly Ser Leu Thr Asp Glu Gln Ile Gln	
-5 1 5	
aaa agg aac aag atc agc aaa gag tgc cag cag gtg tcc gga gtg	231
Lys Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val	
10 15 20	
tcc caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gac	276
Ser Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp	
25 30 35	
gat ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag aaa act	321
Asp Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr	
40 45 50	
gga gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa	366
Gly Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys	
55 60 65	
gcc aag ctg aag cat gtg gcc agc gac gaa gag gtg gac aag atc	411

Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile
70 75 80

gtg cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct 456
Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala
85 90 95

tat gac acc ttc aag tgt att tac gac agt aaa cct gat ttc tct 501
Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser
100 105 110

cct att gat taa ctcgagcacc accaccacca ccactgagat 543
Pro Ile Asp
115

<210> 39
<211> 149
<212> PRT
<213> Tenebrio molitor

<223> Mature protein with His-tag, Clone 7.5

<400> 39
Met Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro
-30 -25 -20

Arg Gly Ser His Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
-15 -10 -5

Gly Ser Leu Thr Asp Glu Gln Ile Gln Lys Arg Asn Lys Ile Ser Lys
1 5 10

Glu Cys Gln Gln Val Ser Gly Val Ser Gln Glu Thr Ile Asp Lys Val
15 20 25 30

Arg Thr Gly Val Leu Val Asp Asp Pro Lys Met Lys Lys His Val Leu
35 40 45

Cys Phe Ser Lys Lys Thr Gly Val Ala Thr Glu Ala Gly Asp Thr Asn
50 55 60

Val Glu Val Leu Lys Ala Lys Leu Lys His Val Ala Ser Asp Glu Glu
65 70 75

Val Asp Lys Ile Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu
80 85 90

Glu Thr Ala Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp
95 100 105 110

Phe Ser Pro Ile Asp
115

<210> 40
<211> 24

<212> DNA
 <213> Tenebrio molitor

 <223> Tm 12.84 upper primer with Bam-H1 site (Clones 2.2, 2.3, 3.4, 3.9, and 7.5)

 <400> 40
 cgcggtatccc tcaccgacga acag 24

 <210> 41
 <211> 25
 <212> DNA
 <213> Tenebrio molitor

 <223> Tm 12.84 lower primer with Xho1 site (Clones 2.2, 2.3, 3.4, 3.9, and 7.5)

 <400> 41
 gagaggataa ctaattgagc tcgcc 25

 <210> 42
 <211> 24
 <212> DNA
 <213> Tenebrio molitor

 <223> Tm 13.17 upper primer with Bam-H1 site

 <400> 42
 cgcggtatccc tgaccgaggc acaa 24

 <210> 43
 <211> 25
 <212> DNA
 <213> Tenebrio molitor

 <223> Tm 13.17 lower primer with Xho1 site

 <400> 43
 gagggtgtaa ctaactgagc tcgcc 25

 <210> 44
 <211> 481
 <212> DNA
 <213> Tenebrio molitor

 <220>
 <221> misc_feature
 <222>
 <223> Consensus of the Tm 12.84 Isoforms, 'n' defined as any nucleotide, 'Xaa' defined as any amino acid.

 <400> 44
 ggcacgagca aaa atg aaa ctc ctc ttg tgc ttt gcn ttc gcc gcc 46

Met Lys Leu Leu Leu Cys Phe Ala Phe Ala Ala
 -15 -10

atc gtc atc gga gct cag gct ctc acc gay gaa cag ata cag aaa 91
 Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys
 -5 1 5

agg aac aag atc agc aaa gar tgc cag cag gng tcc gga gtg tcc 136
 Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Xaa Ser Gly Val Ser
 10 15 20

caa gag acg atc gac aaa gtc cgc aca ggt gtc ttg gtc gay gat 181
 Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp
 25 30 35

ccc aaa atg aag aag cac gtc ctc tgc ttc tcg aag ara act gga 226
 Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly
 40 45 50

gtg gca acc gaa gcc gga gac acc aat gtg gag gta ctc aaa gcc 271
 Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala
 55 60 65

aag ctg aag cat gtg gcc agc gac gaa gar gtg gac aag atc gtg 316
 Lys Leu Lys His Val Ala Ser Asp Glu Glu Val Asp Lys Ile Val
 70 75 80

cag aag tgc gtg gtc aag aag gcc aca cca gag gaa acg gct tat 361
 Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala Tyr
 85 90 95

gac acc ttc aag nnt att tac gac agt aaa cct gat ttc tct cct 406
 Asp Thr Phe Lys Xaa Ile Tyr Asp Ser Lys Pro Asp Phe Ser Pro
 100 105 110

att gat taa ttgttttgta tttgrctgaa ttttgacaat aaaggntanta 455
 Ile Asp
 115

tcgttatgna aaaaaaaaaa aaaaaa 481

<210> 45
 <211> 484
 <212> DNA
 <213> Tenebrio molitor

<220>
 <221> misc_feature
 <222>
 <223> Consensus of Seq ID #44 with Tm 13.17, 'n' defined as any nucleotide,
 'Xaa' defined as any amino acid

<400> 45
 ggcanrnnnnn aar atg aar ytn ctc tnn tgy ytn ryn tyc nyc ryy 46
 Met Lys Leu Leu Xaa Cys Phe Ala Phe Ala Ala
 -15 -10

ntn ntn rtc rna gyt cag gcy ctn acc gan gna car atn nag aaa	91
Xaa Xaa Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys	
-5 1 5	
nng aac aag atc agc aaa rar tgy car nan gnr nny gga gtg tcn	136
Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val Ser	
10 15 20	
caa gag ayn atn rnc aaa gyy cgc ann ggt gnc tng gnn gay gat	181
Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp	
25 30 35	
ccy aaa ntg aar nrn can gty yty tgc ntn ncn arg arn rcy ggn	226
Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly	
40 45 50	
ntg gcn acn gaa ncn gga gan ryn rnn gtn gan gtr ytn arr gnn	271
Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala	
55 60 65	
aag ntg arg nan gtn rcy rrc aac gac gaa gar ryn gan aar atc	316
Lys Leu Lys His Val Ala Ser Asn Asp Glu Glu Val Asp Lys Ile	
70 75 80	
rtn nan aag tgc gyn gtc aag arr gny acn nyn gar gar acg gyn	361
Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala	
85 90 95	
tny ray acy ttc aar nnt rty nnn ran ary aar ccn ran ttc tcn	406
Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser	
100 105 110	
ccn rtt gat tra nynnnyynna ytnngnnnrnr nttyranaat aaagnnnntn	458
Pro Ile Asp	
115	
tnrtnnnrna aaaaaaaaaa aaaaaa	484

<210> 46
 <211> 484
 <212> DNA
 <213> Tenebrio molitor

<220>
 <221> misc_feature
 <222>
 <223> Consensus of Seq ID #45 with B1/B2, 'n' defined as any nucleotide, 'Xaa'
 defined as any amino acid,

<400> 46	
ggcanrnnnn aar atg aar ytn ctc tnn tgy ytn ryn tyy nyc ryy	46
Met Lys Leu Leu Leu Cys Phe Ala Phe Ala Ala	
-15 -10	

ntn ntn rtc nna gyt cag gcy ntn acy nan gna nan ntn nag nna Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys -5 1 5	91
nng nnc nar ayc agc rna rar tgy nar nnn gnr nny gga gtg tcn Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Val Ser Gly Val Ser 10 15 20	136
naa gan ryn atn rnn ara gyy cgc ann ggt gnc tng gnn gay gay Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp 25 30 35	181
ccy aaa ntg aar nnn can nty yty tgc ntn nyn arg rnn nyy grn Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly 40 45 50	226
ntr gyn rcn gaa ncn gga gan ryn rnn gyn gan ryr ytn arr gnn Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala 55 60 65	271
aag ntg ang nrr nnn nnn rnn ann rnn rar rar ryn rrr arr ntn Lys Leu Lys His Val Ala Ser Asn Asp Glu Glu Val Asp Lys Ile 70 75 80	316
nyn nrr arn nnn nnn nnn nng arn rnn nyn nnn rar rnr nnn nnn Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala 85 90 95	361
tnn ran nyn yyn aan nnn nny nnn rrr ann arn ccn rnn tyy tyn Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser 100 105 110	406
cnn ryt rnt trn nnnnnnnnnn ynnngnnrrr nttyranaat aaagnnnytn Pro Ile Asp 115	458
tnrtnnrrna aaaaaaaaaa aaaaaa	484

<210> 47
 <211> 484
 <212> DNA
 <213> Tenebrio molitor

<220>
 <221> misc_feature
 <222>
 <223> Consensus of SEQ. ID #46 with AFP-3, 'n' defined as any nucleotide,
 'Xaa' defined as any amino acid

<400> 47
 ggcnnrnnnn aar atg aar ytn ctc ynn tgy ytn ryn yyy nyy ryy
 Met Lys Leu Leu Cys Phe Ala Phe Ala Ala
 -15 -10

ntn ntn ryc nrr ryy yan gcy ntn acy nan rna nnn nnn nag nrr	91
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Ile Val Ile Gly Ala Gln Ala Leu Thr Asp Glu Gln Ile Gln Lys	
-5 1 5	
nng nny nar nnc agc rnn rnn tgy nar nnn gnr nny gga gtr tcn	136
Arg Asn Lys Ile Ser Lys Glu Cys Gln Gln Glu Ser Gly Val Ser	
10 15 20	
naa gan nyn ntn rnn arr gyy cgc ann ngd gnn nnr gnn gay gay	181
Gln Glu Thr Ile Asp Lys Val Arg Thr Gly Val Leu Val Asp Asp	
25 30 35	
ccy aaa ntg aar nnn can nyy yty tgc ntn nyn arg rnn nyy grn	226
Pro Lys Met Lys Lys His Val Leu Cys Phe Ser Lys Lys Thr Gly	
40 45 50	
ntn ryn rnn gnn nnn ggn gan nyn nnn nyn gan nnn ntn arr rnn	271
Val Ala Thr Glu Ala Gly Asp Thr Asn Val Glu Val Leu Lys Ala	
55 60 65	
aar ntn ang nnn nnn nnn rnn rnn nnn rar rar ryn rrn rrn ntn	316
Lys Leu Lys His Val Ala Ser Asn Asp Glu Glu Val Asp Lys Ile	
70 75 80	
nyn nnn arn nnn nnn nnn nng arn rnn nyn nnn nar nnn nnn nnn	361
Val Gln Lys Cys Val Val Lys Lys Ala Thr Pro Glu Glu Thr Ala	
85 90 95	
nnn ran nyn yyn aan nnn nny nnn rrn ann arn ycn nnn tnn nnn	406
Tyr Asp Thr Phe Lys Cys Ile Tyr Asp Ser Lys Pro Asp Phe Ser	
100 105 110	
cnn nyn rnn trn nnnnnnnnnn ynnnnnnnnn nnnnnnnaat aaannnnnnn	458
Pro Ile Asp	
115	
nnnnnnnnna aaaaaaaaaa aaaaaa	484

<210> 48
 <211> 136
 <212> PRT
 <213> Tenebrio molitor

<220>
 <221> misc_feature
 <222>
 <223> General Consensus of Clones, B1, B2 and AFP-3, 'n' defined as any nucleotide, 'Xaa' defined as any amino acid

<400> 48
 Met Lys Leu Leu Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 -15 -10 -5
 Xaa Ala Xaa Thr Xaa Xaa Xaa Xaa Glx Xaa Xaa Xaa Xaa Xaa Ser Xaa
 1 5 10

Xaa	Cys	Xaa	Xaa	Xaa	Ser	Gly	Xaa	Ser	Glx	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	15	20	25	30
Arg	Xaa	Xaa	Xaa	Xaa	Xaa	Asp	Asp	Pro	Lys	Xaa	Lys	Xaa	Xaa	Xaa	Xaa	35	40	45	
Cys	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Gly	Xaa	Xaa	Xaa	50	55	60	
Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Lys	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Glu	65	70	75	
Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Lys	Cys	Xaa	Val	Xaa	Xaa	Xaa	Thr	Xaa	80	85	90	
Glx	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Lys	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	95	100	105	110
Xaa	Phe	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa									115			